

PCT

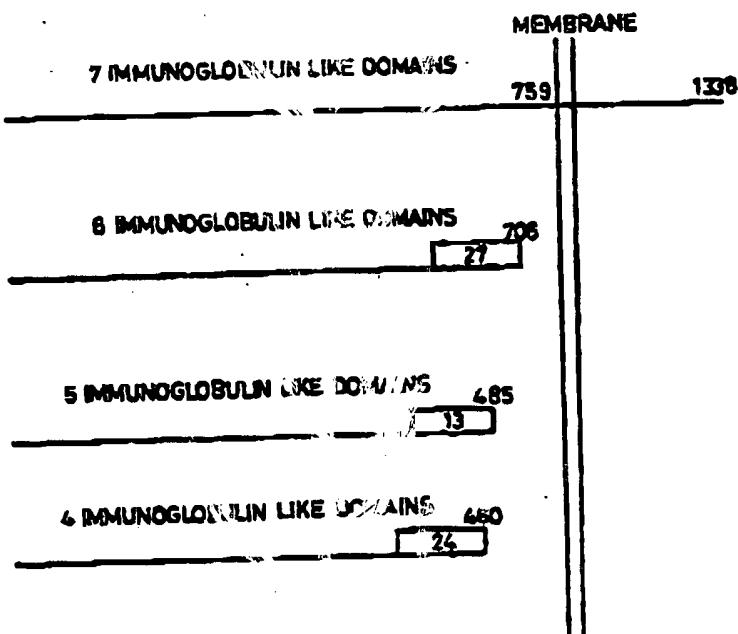
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(54) Title: **FLT-1 (FMS-LIKE TYROSINE-KINASE), FLT-1, VARIANTS THEREOF USES AS GROWTH FACTOR INHIBITORS**



(57) Abstract

Disclosure is an altered, soluble form of the FLT polypeptide being capable of binding to VEGF and thereby exerting an inhibitory effect thereon, the polypeptide comprising five or fewer complete immunoglobulin-like domains, together with pharmaceutical compositions comprising the polypeptide, and various uses thereof.

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FLT-4 (fms-like Tyrosine kinase), FLT-15, variants thereof used as growth factor inhibitors

Field of the Invention

This invention relates to substances which inhibit growth factors, in particular, vascular endothelial growth factor (VEGF), methods of inhibiting growth factors and of treating tumours and regulating fertility.

Background of the Invention

A considerable number of human growth factors are now known, many of which have been at least partly characterised. Among them is vascular endothelial growth factor (VEGF), which has been identified in several tissues (Gospodarowicz *et al.*, 1989 PNAS 86, 7311-7315; Conn *et al.*, 1990 PNAS 87, 2628-2632; Tischer *et al.*, 1991 J. Biol. Chem. 266, 11947-11954). As its name suggests, this growth factor is a highly specific mitogen for endothelial cells and is greatly involved in angiogenesis. VEGF is a homodimeric glycoprotein of two 23kDa subunits exhibiting sequence homology with platelet-derived growth factor A and B chains and placenta growth factor.

The homologous tyrosine kinase receptors fms-like tyrosine kinase receptor (FLT) and kinase insert domain-containing receptor (KDR) function as high-affinity VEGF receptors (de Vries *et al.*, 1992 Science 255, 989-991; Terman *et al.*, 1992 Biochem. Biophys. Res. Commun. 187, 1579-1586). Both FLT and KDR are membrane-spanning receptors that each contain seven immunoglobulin-like domains in the extracellular ligand-binding region, an intracellular tyrosine kinase domain and a transmembrane domain. The transmembrane domain serves to anchor the receptor in the cell membrane of the cells in which it is expressed.

A number of membrane-bound receptor molecules have been found to exist in truncated soluble forms, generated either by proteolytic processing or by alternative splicing of

mRNA. Recently, Kendall & Thomas (1992 PNAS 90, 10,705-10,709, and WO94/21679) described the discovery of a soluble form of FLT receptor (sFLT) generated by alternative splicing.

Essentially, Kendall & Thomas screened a human umbilical vein endothelial cell (HUVEC) cDNA library with one probe specific for the 3' end of the flt coding region (encoding the intracellular tyrosine kinase domain) and with another probe specific for the 5' flt coding portion (encoding one of the extracellular N terminal domains). Clones which hybridised with the 5' specific probe but not with the 3' specific probe were selected for further study. In this way, a clone was isolated which encoded a soluble FLT polypeptide lacking the transmembrane domain and the intracellular domain. The truncation resulted from "readthrough" to an intronic termination codon. It was suggested by Kendall & Thomas that the soluble receptor could act as an efficient specific antagonist of VEGF *in vivo*.

The present invention is based on the discovery of further soluble variants of FLT, the existence of which was not predicted by the teaching of Kendall & Thomas.

Summary of the Invention

In a first aspect the invention provides an altered, soluble form of the FLT polypeptide being capable of binding to VEGF and thereby exerting an inhibitory effect thereon, the polypeptide comprising five or fewer complete immunoglobulin-like domains. Preferably, the altered FLT polypeptide comprises four or fewer complete Ig-like domains. The altered soluble FLT polypeptide inhibits VEGF by preventing it binding to its natural receptors, flt and KDR, present on the surface of target cells. Surprisingly, such truncated forms, lacking a major extracellular portion of the molecule, are believed to retain affinity for VEGF.

The term "soluble" as used herein is intended to refer to altered forms of the FLT polypeptide which do not comprise a transmembrane domain and thus generally do not become associated with the cell membrane of cells in which the molecule is expressed. In particular, the invention provides soluble altered forms of the FLT polypeptide

consisting substantially of four or five complete immunoglobulin-like domains.

In a particular embodiment the invention provides an altered, soluble form of FLT having at its C-terminus a region substantially having the amino acid sequence of the sequences termed FLT4 or FLT15 shown in Figure 5, or a functional equivalent thereof. The term "functional equivalent" as used above is intended to include those polypeptides which have substantially the same deletions as the polypeptides encoded by FLT4 or FLT15 (with respect to the unaltered full length FLT molecule), but which may also have other deletions, additions or substitutions, (in particular conservative substitutions), and which retain an inhibitory effect for VEGF.

Preferably the polypeptide will also comprise, at its N-terminus, the amino acid sequence substantially corresponding to the equivalent portion of the unaltered wild-type FLT polypeptide. Conveniently, polypeptides in accordance with the invention will comprise around 400 to 500 amino acid residues, preferably around 480 amino acid residues, most preferably between 480 and 440 amino acid residues of the wild type FLT sequence.

Preferably the polypeptides of the invention arise by alternative splicing of mRNA or by proteolytic processing of a mature polypeptide, although it will be apparent to those skilled in the art that the polypeptide could be encoded by a nucleic acid derived, at least in part, by recombinant DNA technology.

In a further aspect the invention provides a nucleic acid sequence encoding a polypeptide in accordance with the invention. In a particular embodiment the invention provides a nucleic acid comprising the sequence of nucleotides inserted at position 1655 of the FLT 4 sequence shown in Figure 3 or the sequence of nucleotides inserted at position 1555 of the FLT 15 sequence shown in Figure 3, or a functional equivalent thereof. Examples of functionally equivalent nucleic acids include those sequences which encode substantially the same polypeptide as those encoded by FLT4 or FLT15 but which differ in nucleotide sequence as a result of the degeneracy of the genetic code. It will be apparent to those skilled in the art that the portion of the inserted nucleotide sequence in FLT4 and FLT15 occurring after the premature termination codon could be omitted without affecting the characteristics of the encoded polypeptide. Accordingly, nucleic acid molecules without

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such sequences are also regarded as functionally equivalent for the purposes of the present invention.

Conveniently, the nucleic acid will substantially comprise the nucleotide sequence of FLT4 or FLT15 shown in Figure 3, together with the nucleotide sequence encoding the N-terminus of unaltered, wild-type FLT. Advantageously, the nucleic acid will be obtainable by means of PCR amplification from a sample of human cells. Desirably, the nucleic acid will be obtainable by means of PCR using primers intended to hybridise to non-conserved regions of the FLT molecule. Conveniently, the nucleic acid sequence will be obtainable by use of PCR primers designed to hybridise to the regions of the FLT sequence shown underlined in Figure 3, or immediately adjacent thereto. In particular, the PCR primers will conveniently have substantially the sequence: 5'- GCAAGGTGTGACTTTGTTTC -3' and 5' - AGGATTCTTCCCTGTGTA -3'.

In another aspect, the invention provides a method of inhibiting VEGF *in vitro*, comprising adding an effective amount of the polypeptide defined above. It may also be desirable to inhibit VEGF in a human subject. Thus the invention provides a method of inhibiting VEGF in a human subject, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance. In particular, VEGF provides a mitogenic stimulus (particularly involved in angiogenesis), so inhibition of VEGF would be expected to provide therapeutic effects in the treatment of tumours or disorders involving inappropriate neovascularisation.

In particular the invention provides for a method of treating tumours or diseases involving inappropriate neovascularisation, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance. Suitable diseases which might be amenable to treatment include ovarian cancer and ovarian hyperstimulation (Boocock *et al.*, 1995 *J. Natl. Cancer Inst.* 87, 506-516).

Furthermore, it has been conclusively demonstrated that FLT is expressed by trophoblasts and cells from ovarian and endometrial tissues (Charnock-Jones *et al.*, 1994 *Biology of Reproduction* 51, 524-530), which clearly suggests a role for VEGF in the growth and

differentiation of trophoblasts during implantation.

Thus, in particular, the invention provides a method of affecting the growth and/or migration of trophoblasts, ovarian or endometrial cells by inhibiting the action of VEGF, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance.

It will be appreciated by those skilled in the art that the identification of FLT on the surface of trophoblasts and endometrial cells also provides a number of possible methods of regulating fertility. For example, the growth of trophoblasts is essential for successful implantation of the embryo. Inhibition of trophoblast growth thus provides a method of contraception or contragestion.

Thus in a further aspect the invention provides a method of regulating the fertility of a human female, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance. An "effective amount" of the polypeptide is an amount sufficient to substantially block the stimulus of VEGF on trophoblasts and/or endometrial cells. Typically, the method will result in reducing the fertility of the female.

Moreover, it might be possible to identify agents which can enhance the effect of VEGF on trophoblasts, and thereby improve the probability of successful implantation, either in assisted or spontaneous cycles. Candidates for such VEGF-enhancing agents would include anti-sense equivalents of the nucleic acid sequences encoding the truncated FLT polypeptides of the invention. It will be apparent to those skilled in the art that these could be used to improve the fertility of a human female.

In a further aspect the invention provides a pharmaceutical composition comprising the polypeptide defined above, together with a physiologically acceptable carrier substance. The composition could be used *in vivo* any one of the methods defined above. In yet another aspect the invention provides for the use of a polypeptide in accordance with the invention in the preparation of a therapeutic composition for the treatment of tumours and

diseases involving inappropriate neovascularisation. Examples of such conditions and diseases are detailed, inter alia, in WO94/10202 and WO94/21679. The invention also includes within its scope a method of making a pharmaceutical composition, comprising mixing the polypeptide defined above together with a physiologically acceptable carrier substance.

The invention will now be described by way of the following illustrative examples and with reference to the drawings, of which:

Figure 1 shows an amino acid multiple alignment of closely related tyrosine kinase receptors (flt, fms and kit, "kit" being another name for KDR);

Figure 2 shows typical results of agarose gel electrophoresis demonstrating the existence of alternatively-spliced flt-coding sequences in various tissue samples;

Figure 3 shows the nucleotide sequence of the 3' region of the sequences encoding full length VEGF receptors (FLT and the related receptor KDR), together with two sequences, FLT4 and FLT15, which encode polypeptides according to the invention;

Figure 4 is a schematic representation of wild type and mutant FLT molecules; and

Figure 5 shows the C terminal amino acid sequences of two polypeptides in accordance with the invention.

Example

Expression of FLT, the VEGF receptor, was investigated in cell lines derived from human trophoblast-like and ovarian and endometrial carcinomas. The trophoblast-like (choriocarcinoma) cell line used was BeWo (obtained from the American Type Culture Collection, Rockville MD, USA). The endometrial carcinoma cell lines were Ishikawa (obtained from Professor M Nishide, University of Tsukuba, Japan), and HEC 1-A and HEC 1-B (from ATCC, USA). The ovarian cancer cell lines were 7, 17R, 25, 25R and 35. These were all shown to be of epithelial origin and had been established in culture

for 10-30 passages. Cell lines 17R and 25R were derived after chemotherapy and subsequent relapse (line 25R originating from the same patient as line 25).

BeWo cells were grown in Ham's F12, according to ATCC recommendations. Endometrial carcinoma lines were grown in McCoy's medium (ICN Flow Laboratories, Irvine, UK) with 10% foetal calf serum (ICN Flow) plus 2mM L-glutamine (ICN Flow) and 50U/ml and 50mg/ml penicillin/streptomycin (ICN Flow).

It was decided to investigate expression of FLT in these cell lines and normal tissues by performing PCT and *in situ* hybridization. It was therefore necessary to construct suitable oligonucleotide primers and probes.

To help design appropriate primers, a protein multiple alignment of closely related tyrosine kinase receptors (FLT, FMS and KIT) was constructed (shown in Figure 1) using the computer program "pileup". This revealed regions of divergent sequence among this family of receptors. The regions chosen for primer design are shown with double underlining in Figure 1. The following nested PCR primers were then synthesized based on these protein sequences:

- A) 5' GCAAGGTGTGACTTTTGTC 3'
- B) 5' GCGCTCGAGAGCATCACTCAG 3'
- C) 5' GCGCGGCCGCAGTAAAATCCA 3'
- D) 5' AGGAATTCTTCCCCTGTGTA 3'

The underlined portions of these oligonucleotides are the regions which hybridise to the *fit* cDNA sequence. The other nucleotides were added to facilitate directional cloning. The cycles used were: first round with primers A and D [95°C 30 seconds, 55°C 30 seconds, 72°C 30 seconds] x 25; second round with primers B and C: [95°C 30 seconds, 44°C 30 seconds, 72°C 30 seconds] x 2 [95°C 30 seconds, 65°C 30 seconds, 72°C 30 seconds] x 25. The internal primers B and C had sites for the restriction enzymes Xba I and Eco R I respectively at their 5' ends to permit directional cloning of the products.

It was found that certain tissues gave rise to PCR amplification products of notably larger size (as judged by agarose gel electrophoresis) than observed for the full length FLT cDNA product. Typical results are shown in Figure 2.

PCR products obtained using the nested set of primers A-D were run out on a gel. Lanes 1-3 are products obtained from primary tissue samples of the ovarian carcinomas designated 17, 17R and 25R. Lanes 4 to 7 are products obtained from cell lines established from the ovarian carcinomas 7, 17R, 25 and 25R. Lanes 8 to 10 are the cell lines HEC 1-A, HEC 1-B and Ishikawa respectively. Lane 11 contains products from HUVECs.

The standard size band was of the expected size (around 285bp) and was found to be identical to the 3' end of the published *fit* sequence (Shibuya *et al.*, 1990 *Oncogene* 5, 519-524). However it can be clearly seen that in addition to the full length *fit* cDNA PCR-amplified product, in lanes 2 (17R, primary tissue) and 4 (7, cell line) are larger bands of approximately 360bp. A faint band of similar size was also apparent in lane 5 (17R, cell line) but is not clearly seen when the gel photograph is reproduced. These larger bands were extracted from the gel by known techniques and subcloned into the plasmid vector pBluescript II KS and then subjected to sequence analysis using the dideoxynucleotide sequencing method (Sanger *et al.*, 1977 *PNAS* 71, 5463-5467).

Sequencing of five independent clones (Boocock *et al.*, 1995 *J. Natl. Cancer Inst.* 87, 506-516) revealed that each contained one of two novel insertions within the published *fit* sequence, in the region between the primers. Three of these clones (termed *FLT5*, *FLT15* and *FLT16*) contained an 85bp insertion at about position 1555, whilst two other clones (*FLT13* & *FLT14*) contained a 65bp insertion at about position 1665 (see Figure 3, numbering based on that of Shibuya *et al.*, 1990 cited above). The insertions account for the larger band size of the PCR products. However, both insertions contain an in-frame termination codon, so that corresponding full length RNAs would encode soluble, truncated receptor variants comprising the first five immunoglobulin-like domains of the extracellular region, up to amino acid 517 or 553, with either 24 or 14 (of which 13 are additional) unrelated amino acids at the C-terminus.

Although these variant fit clones were derived from partial cDNAs encoding only amino acids 503 onward, PCR products of the sizes predicted for corresponding full length cDNAs were amplified from cDNA derived from HUVEC cells, human chorion and ovarian carcinoma cell line 7, using primers specific for each of the novel insertions together with a primer binding just 5' of the initiating ATG (data not shown).

Figure 4 is a schematic representation of various FLT receptor molecules. At the top, (a) shows the wild type, full length FLT receptor molecule, (b) represents the truncated version described by Kendall & Thomas, (c) represents the polypeptide encoded by FLT4 and (d) represents the polypeptide encoded by FLT15. The numerals at the right show the number of amino acids in the molecule and numerals in the boxes represent the number of amino acids present in the sFLT variants but not in the wild type molecule.

Figure 5 shows the predicted C terminal amino acid sequence of the polypeptides which would be encoded by "full length" FLT4 and FLT15 clones (i.e. clones which contained all the nucleotide sequence 5' of the primer site used to generate the actual clones). The last 14 amino acids of the FLT4 clone, and the last 24 amino acids of the FLT15 clone, are divergent from the wild type FLT sequence.

Claims

1. An altered, soluble form of the FLT polypeptide being capable of binding to VEGF and thereby exerting an inhibitory effect thereon, the polypeptide comprising five or fewer complete immunoglobulin-like domains.
2. A polypeptide according to claim 1, comprising four or fewer complete immunoglobulin-like domains.
3. A polypeptide according to claim 1 or 2, having at its C terminus substantially the amino acid sequence of FLT4 as shown in Figure 5, or a functional equivalent thereof.
4. A polypeptide according to claim 1 or 2, having at its C terminus substantially the amino acid sequence of FLT15 as shown in Figure 5, or a functional equivalent thereof.
5. A polypeptide according to any one of the preceding claims, comprising around 400 to 500 amino acid residues of the wild type FLT polypeptide.
6. A nucleic acid sequence encoding a polypeptide in accordance with any one of the preceding claims.
7. A nucleic acid sequence according to claim 6, comprising the sequence of the nucleotides inserted at position 1655 of the FLT4 sequence shown in Figure 3, or a functional equivalent thereof.
8. A nucleic acid sequence according to claim 6, comprising the sequence of the nucleotides inserted at position 1555 of the FLT15 sequence shown in Figure 5, or a functional equivalent thereof.
9. A method of inhibiting VEGF *in vivo*, comprising adding an effective amount of a polypeptide in accordance with any one of claims 1 to 5.

10. A method of inhibiting VEGF in a human subject, comprising administering an effective amount of a polypeptide in accordance with any one of claims 1 to 5, together with a physiologically acceptable carrier substance.
11. A method according to claim 10, comprising the use of a polypeptide in accordance with any one of claims 1 to 5 in the treatment of tumours or diseases involving inappropriate neovascularisation.
12. A method according to claim 11, for the treatment of ovarian cancer, ovarian hyperstimulation, or endometriosis.
13. A method of affecting the growth and/or migration of trophoblasts, ovarian or endometrial cells by inhibiting the action of VEGF by administration of an effective amount of a polypeptide in accordance with any one of claims 1 to 5, together with a physiologically acceptable carrier substance.
14. A method of regulating the fertility of a human female by administration of an effective amount of a polypeptide in accordance with any one of claims 1 to 5, together with a physiologically acceptable carrier substance.
15. A pharmaceutical composition for use in the method of any one of claims 11 to 14, comprising a polypeptide in accordance with any one of claims 1 to 5, and a physiologically acceptable carrier substance.
16. A method of making a composition according to claim 15, comprising mixing a physiologically acceptable carrier substance together with a polypeptide according to any one of claims 1 to 5.

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kit MRGARGAWDF LCVLLLLLRLV QTGSSQPSVS PGEPSPPSIH PGKSDLIVRV
 fmsM GPGVLLLLLV ATAWHGQGIP VIEPSVP... .ELVVKP
 flt MVSYWDITGVL ICAILSCLL TGSSSGSKLK DPELSLKG... .TOHIMOA
 51
 kit GDEIRLLCTD PGFVKW.... .TEIILD ETENENQ... .NEWITE
 fms GATVTILRCVG NGSVEWGDGP.ASPHWT LYSDGSS... .SILSTN
 flt GOTLHLQCRG EAAHKWSLPE MVSKESERLS ITKSACGRNG KOFCASTLTIN
 101
 kit KAEATNTGKY TC....T... NKGGLNSIY VFVRDPAKLFLVDRS
 fms NATFQNTGTY RC....TEPG DPLGGSAAIH LYVKDPARPWNVLAQE
 flt TAQANHTGFY SCKYLAVPTS KKETESAIY IFISDTGRPF VEMYSEIPEI
 151
 kit LYKGEDNDTL VRCP LTDPEV .TNYSLKGCO GPLPKD.LR FIPDPKALIM
 fms VVFEDQDAL LPCLLTDPV L EAGVSLRVR GRPLMRH.TN YSFSPWHGFT
 flt IHMTEGRELV IPCRVTSRNI ..TVTLKKFP LDTLIPDGKR IIWDSRKGFI
 201
 kit IKSVKRAYHR LCLHCSVDQE GKSVLSEKFI LKVRPAFKAV PWSVSKASY
 fms IHRAK.FIQS QDYQCSALMG GRKVMSISIR LKVQKVIPGP PALTLPFEL
 flt ISNAT.YKEI GLITCEATVN GLYKTNYL T HQTNTIIDV QISTPRP'KL
 251
 kit LLREGEETV TCTI.KDVSS SVYSTWKREN SQTKLOEK... .YNSWHHG
 fms VRIRGEAAQI VCSA.SSVDV NFDVFLQHNN ..TKLAIP... .QQSDFHNN
 flt L..RGHTLVL NCTATTPLNT RVQMTWSYPD EKNKRASVRR RIDQSNSHAN
 301
 kit FNYERQATLT ISSARVNDSG VEMCYANNTE GSANVTTLE VWDKGFINI.
 fms .RYQKVLTIN LDQVDFQHAG NYSCVASNVO GHHSSTMFR VVESAYINL.
 flt IFYS...VLT IDKMQNKDKG LYTCRVRSGP SFKSVNTSVH IYDKAFITVK
 351
 kit FPMINTTVFV NDGENVDLIV EYEAFPKPEH QOWIYMNRTF TDKWEDYAPKS
 fms SSEQNLIQEV TVGEGLNLIK MVEAYPGLQGFNWTY LGPFSDHQPE
 flt HRKQOVLETV AGKRSYRLSM KVKAEPSPREV V..... .WLKD
 401
 kit ENESN..... IRYVSELHL TRLKGTEGGT YTFLVS..NS DVNAALIAFN
 fms PKLANATTKD TYRHFTLSSL PRKPKSEAGR YSFALAR..NP GGWRALTFEL
 flt GLPATEKSAR YLTRGYSLII KDVTEEDAGN YTILLSIKOS NVENLNLATL

Fig. 1 Sheet 1

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	451	500
kit	YVNTKPEI.. LTYDRL.... VN.. GML QCVAAFGPEP TIDWYFCPGT	
fms	TLRYPPEV.. SVIWF.... INSGSTL LCAASGYPQP NVIWLOC&GH	
flc	IVNVKPOIYE KAVSSFPDPA LYPLGSRQIL TCTAYGIPQP TIKWEWHPCN	
	501	550
kit	EQRC.....	
fms	TDRCD.....	
flc	<u>HNHSEARCOF CSNNEESFIL DADSNMGNRI ESITORMAI</u> I EGKNKMASTIL	
	551	600
kit	SASV
fms	EAV
flc	VVADSRISGI YICIASNKVG TVGRNISFYI TDVPNFGHVN LEKMPTE <u>ED</u>	
	601	650
kit	LPV.. DVQTL NSSGPPE...	GLVVQSS
fms	LQWDDDPYPE VLSQEPF...	HKVTVQSL
flc	LKLSCTVNKF LYRDVTWILL RTVNNRIMHY SISKOKMAIT KEHSITLNAT	
	651	700
kit	IDSSAFKHNG TVECKAYNDV G.....	
fms	LTVETLEHNO TYECAHNSV G.....	
flc	IMNVSLQDSG TYACRAPNVY <u>TGEEILOKKE ITIRDQEAPY LLRNLSDHIV</u>	
	701	750
kit	..KTSAYFNE A..... FKGNKNEQ IHPHTLFTP.	
fms	..SGSWAF. I P..... ISAGAHTH PPDEFILFTP.	
flc	AISSSTTLDC HANGPEPOQI 1WFKNHHKIQ QEPGIILGPG SSTLFIEFVT	
	751	800
kit	LLI GEVTIVAGMC
fms	WV ACMSIMALL
flc	EEDEGVYHCK ATNQKGSVES SAYIVQTS DKSNELITL TCIC/AATLIF	
	801	850
kit	IVMILTYKY LOKPMYEVQW KVVEEINGNN YVYID.. PTQ LPYDH.KMEF	
fms	LLLLLLLYKY KOKPKYQWRW KIIIESYEGNS YTFID.. PTQ LPYNE.KMEF	
flc	WLLLTLLIRK MKRSSSEIKT DYLSIIMDPD EVPLDEQCER LPYDASKMEF	
	851	900
kit	PRNRLSFGKT LGAGAEGKVV EAYAYGLIKS DAAMTVAVKM LKPSAHYTER	
fms	PRNNLOFGKT LGAGAEGKVV EATAEGLGKE DAVLKVAVKM LKSTAHADEK	
flc	ARERLKLGKS LGRAFGKVV QASAFGIIKKs PTCRTVAVKM LKEGATASEY	

Fig. 1 Sheet 2

SUBSTITUTE SHEET (RULE 26)

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	901	950
kit	EALMSELKVL SYLGNHMNIV NLLGACT.IG GPTLVITEYC CYGDILLNFLR	
fms	EALMSELKIM SHLGHENIV NLLGACT.HG GPVLVITEYC CYGDILLNFI.R	
flt	KALMTELKIL THIGHLNWV NLLGACTKQG GPLMVTEYC KYGNLSNYIK	
	951	1000
kit	RKRDSFI....C...SKQE DHAEAAALYKN L.....LHS KESSCSDDSTN	
fms	RKAFAAML...GPSLSPGQ DPEGGVVDYKN IHLEKKYVRR DSGFSSQGVD	
flt	SKRDLFFLNK DAALKMEPKK EKMEPGLEQG KKPRLDSVTS SESFASSGFQ	
	1001	1050
kit	EYMDMKPGVS YVVPPTKADKR RSVRIGSYIE RQVTPAIMED DELAIDLECL	
fms	TYVEMRP....VSTSSN....DSFSE QDLD....KE DGRPIELRCL	
flt	EDKSL.....SDVEE EEDSDGFYKE ...PYTMELL	
	1051	1100
kit	LSEFSYQVAKG MAFIASKNCI HRDLAARNIL LTHGRITKIC DFGLARDIMN	
fms	LHFSSQVAQG MAFIASKNCI HRDVAARNVL LINGHVAKIG DFGLARDIMN	
flt	ISYSFQVARG MEFLSSRKCI HRDLAARNIL LSENNVVKIC DFGLARDIMK	
	1101	1150
kit	DSNYVVKGNA RLPVKWMAPE SIFNCVYTTFE SDWWSYGIFL WEIFSLGSSP	
fms	DSNYIVKGNA RLPVKWMAPE SIFDCVYTVO SDWWSYGILL WEIFSLGLNP	
flt	NPDYVRKGDT RLPLKWMAPF SIFUKIYSTK SDWWSYGVLL WEIFSLGGSP	
	1151	1200
kit	YPGMPVDSKF YKMIKEGFRM LSPEMAPAEM YDIMKTOWDA DPLKREIHKQ	
fms	YPGILVNSKF YKLVKDGYOM AOPAFAPRNI YSIMQACWAL EPTHRPTFOQ	
flt	YPGVQMDDEF CSRLREGMRM RAPEYSTPEI YQIMLDCWHR DPKERPREAE	
	1201	1250
kit	IV....OLIE KQISES.TNH I.....Y SNLANCSPNR QKPVUDH5VR	
fms	IC....SFLO EQAQEDRRER D.....Y TNLPSSSR.S.GGSGS	
flt	LVEKLGDLLQ ANVQODGKDY IPINAILTGN SGFTYSTPAF SEDFFKESIS	
	1251	1300
kit	INSVGSTASS SQP.....L LVHDDV.....	
fms	SSSELEEESS SEH.....L TOLEQGDIAQ PLLOPNNYOF C.....	
flt	APKFNSGSSD DVRYVNAFKF MSLERIKTFE ELLPNATSME DDYQGDSSTL	
	1301	1350
kit	
fms	
flt	LASPMLKRET WTDSKPKASL KIDLRTVTSKS KESGLSDVSR PSFHSSOGH	
	1351	1389
kit	
fms	
flt	VSEGKRRTTY DHAEELERKIA CCSPPPDYNS WLYSTPPI	

Fig.1 Sheet 3

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1 2 3 4 5 6 7 8 9 10 11



Fig. 2

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		1410
KDR	AGAGTGCGCC	AACGAGCCCA GCCAAGCTGT CTCAGTGACA AACCCATACC
FLT	ACCCCTGTAA	CCATAACATT CCGAAGCAAG <u>GATGACTTT</u> TGTCCAATA
	1460	
KDR	CTTGTGAAGA	ATGGAGAAAGT GTGGAGGAAC TCCAGGGAGG AAATAAAATT
FLT	ATGAAGAGTC	CTTTATCCTG GATGCTGACA GCAACATGGG AAACAGAATT
	1510	
KDR	GAAGTTAATA	AAAATCAATT TGCTCTAATT GAAGGAAAAA ATAAA-----
FLT	<u>GAGAGCATCA</u>	<u>CTCAGCGCAT</u> GGCAATAATA GAAGGAAAGA ATAAAG-----
FLT4	<u>GAGAGCATCA</u>	<u>CTCAGCGCAT</u> GGCAATAATA GAAGGAAAGA ATAAAG-----
FLT15	<u>GAGAGCATCA</u>	<u>CTCAGCGCAT</u> GGCAATAATA GAAGGAAAGA ATAAAGCTTCC
KDR	-----	-----
FLT	-----	-----
FLT4	-----	-----
FLT15	ACCAGCTGAC	AGTTCTTCA TGTTGCCACC TACAAGCTTC TTTTCCAACT
	1555	
KDR	-----	CTGTAAGTAC CCTTGTTATC
FLT	-----	ATGGCTAGCA CCTTGTTTGT
FLT4	-----	ATGGCTAGCA CCTTGTTTGT
FLT15	ACTTCCATTT	CCTTCCGTGA CTCTAACCGG ATGGCTAGCA CCTTGTTTGT
	1575	
KDR	CAAGCGGCAA	ATGTGTCAAGC TTGTACAAA TGTGAAGCGG TCAACAAAGT
FLT	GGCTGACTCT	AGAATTTCIG GAATCTACAT TTGCATAGCT TCAATAAAAG
FLT4	GGCTGACTCT	AGAATTTCIG GAATCTACAT TTGCATAGCT TCAATAAAAG
FLT15	GGCTGACTCT	AGAATTTCIG GAATCTACAT TTGCATAGCT TCAATAAAAG
	1625	
KDR	CGGGAGAGGA	GAGAGGGTGA TCTCCTTCCA CGTGACCAGG -----
FLT	TTGGGACTGT	GGGAAGAAAC ATAAGCTTTT ATATCACAGA -----
FLT4	TTGGGACTGT	GGGAAGAAAC ATAAGCTTTT ATATCACAGA ATTGTCAAAC
FLT15	TTGGGACTGT	GGGAAGAAAC ATAAGCTTTT ATATCACAGA -----
KDR	-----	-----
FLT	-----	-----
FLT4	TTTGAGTGCQ	TTGATCCCTG CTCTCAGGAA TAGAACTCTA CTCATCGGA
FLT15	-----	-----

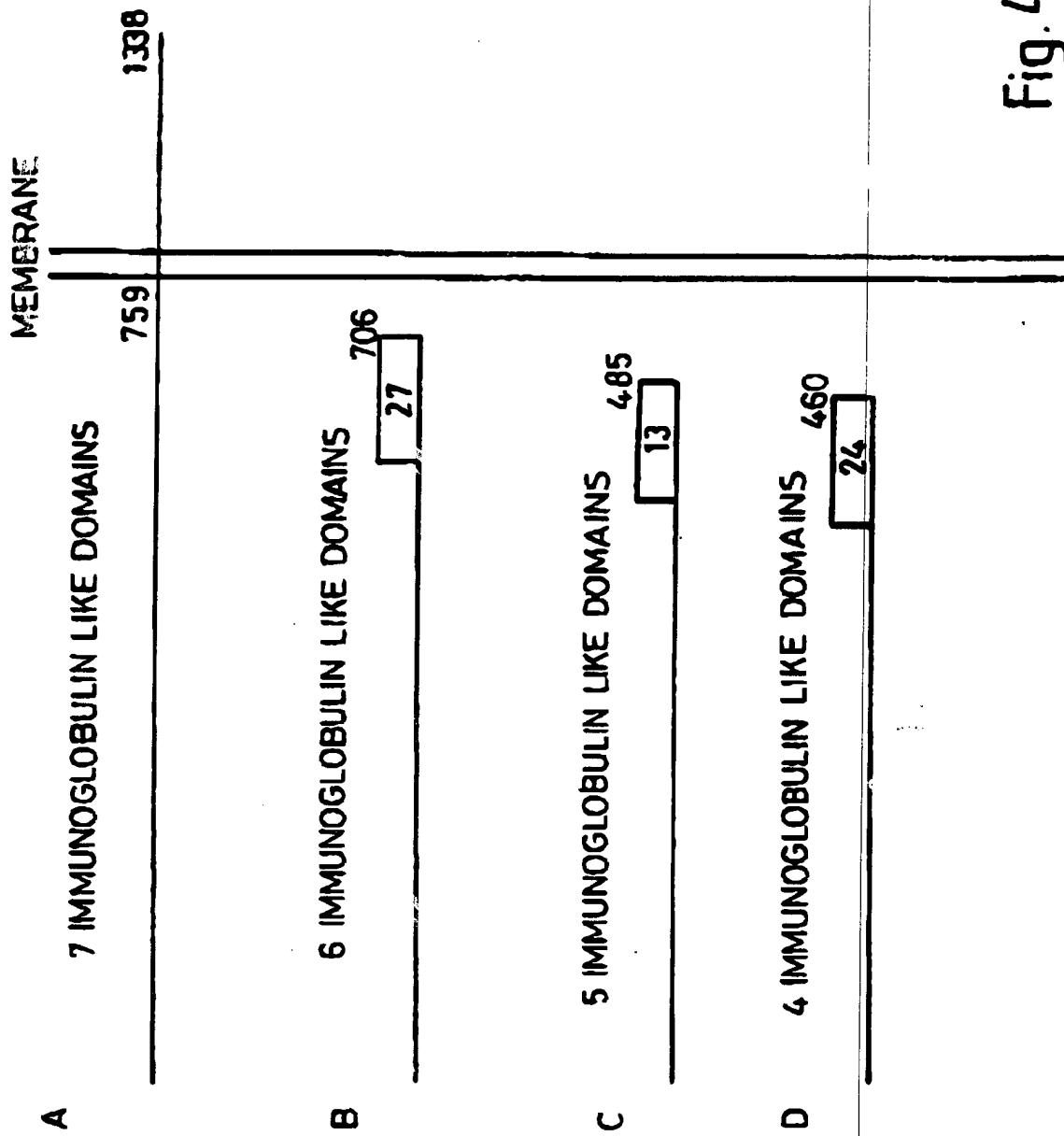
Fig. 3 Sheet 1

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	1665	
KDR	-----GGTCC T---GAAATT ACTTGCAAC CTGACATGCA	GCCCCACTGAG
FLT	-----TGTGC CAAATGGGTT TCATGTTAAC TTGGAAAAAA	TGCCGACCGA
FLT4	TCTCATGTGC CAAATGGGTT TCATGTTAAC TTGGAAAAAA	TGCCGACCGA
FLT15	-----TGTGC CAAATGGGTT TCATGTTAAC TTGGAAAAAA	TGCCGACCGA
	1710	
KDR	CAGGAGAGCG TGTCTTTGIG GTCGCAGTGCA GACAGATCTA	CGTTTGAGAA
FLT	AGGAGAGGAC CTGAAACTGT CTTGCACAGT TAACAAGTTC	TTATACAGAG
FLT4	AGGAGAGGAC CTGAAACTGT CTTGCACAGT TAACAAGTTC	TTATACAGAG
FLT15	AGGAGAGGAC CTGAAACTGT CTTGCACAGT TAACAAGTTC	TTATACAGAG
	1760	
KDR	CCTCACATGG TACAAGCTTG GCCCACAGCC TCTGCCAATC	CATGTGGGAG
FLT	<u>ACGTTACTTG GATTTTACCG CGGACAGTTA ATAACAGAAC</u>	AAATGCACTAC
FLT4	<u>ACGTTACTTG GATTTTACCG CGG</u>	
FLT15	<u>ACGTTACTTG GATTTTACCG CGG</u>	
	1810	
FLT	AGTATTAGCA AGCAAAAAAT GGCCATCACT AAGGAGCACT CCATCACTCT	
	1860	
FLT	TAATCTTACC ATCATGAATG TITCCCTGCA AGATTCAAGGC	AATATGCCT
	1910	
FLT	GCAGAGCCAG GAATGATAAC ACAGGGGAAG AAATCCTCCA	GAAGAAAAGAA

Fig. 3 Sheet 2

Fig. 4



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FLT4

1 ESITQRMAII EGKNKMASTL VVADSRISGI YICIASNKVG TVGRNISFYI
51 TELSNFECLH PCSQE*

FLT15

1 ESITQRMAII EGKNKLPPAD SSFMLPPTSF SSNYFHFLP*

Fig. 5

INTERNATIONAL SEARCH REPORT

International application No.
PCT/GB 95/01213

A. CLASSIFICATION OF SUBJECT MATTER
IPC 6 C12N15/12 C07K14/71 A61K38/17

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
IPC 6 C12N C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	CRIT REV ONCOG, 1993, 4 (6) P595-613, UNITED STATES, ROSNET O ET AL 'Hematopoietic receptors of class III receptor-type tyrosine kinases.' see the whole document ---	1,2,9-16
Y	ONCOGENE, vol. 8, no. 11, November 1993 ENGLAND, pages 2931-2937, PAJUSOLA, K. ET AL.; 'Two human FLT4 receptor tyrosine kinase isoforms with distinct carboxy terminal tails are produced by alternative processing of primary transcripts' see the whole document ---	1,2,9-16

Further documents are listed in the continuation of box C.

Patent family members are listed in annex.

* Special categories of cited documents :

- 'A' document defining the general state of the art which is not considered to be of particular relevance
- 'E' earlier document but published on or after the international filing date
- 'L' document which may throw doubt on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- 'O' document referring to an oral disclosure, use, exhibition or other means
- 'P' document published prior to the international filing date but later than the priority date claimed

- 'T' later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
- 'X' document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
- 'Y' document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.
- 'Z' document member of the same patent family

Date of the actual completion of the international search

17 October 1995

Date of mailing of the international search report

08.11.95

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Authorized officer

Nauche, S

INTERNATIONAL SEARCH REPORT

International application No	
PCT/GB 95/01213	

C(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	ONCOGENE, AUG 1993, 8 (8) P2293-8, ENGLAND, FINNERTY H ET AL 'Molecular cloning of murine FLT and FLT4.' see the whole document ----	1, 2, 9-16
Y	WO,A,94 01576 (SYSTEMIX INC) 20 January 1994 see the whole document ----	1, 2, 9-16
A	WO,A,93 15201 (NEW ENGLAND DEACONESS HOSPITAL) 5 August 1993 see the whole document ----	1-16
A	WO,A,92 14748 (AMERICAN CYANAMID CO) 3 September 1992 see the whole document -----	1-16

INTERNATIONAL SEARCH REPORT

Intern. Appl. No.

PCT/GB95/01213

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. Claims Nos.: 10-13
because they relate to subject matter not required to be searched by this Authority, namely:
Remark: Although claims 10-13 are directed to a method of treatment of the human/animal body as well as diagnostic methods (Rule 39.1(iv) PCT) the search has been carried out and based on the alleged effects of the compound/composition.
2. Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remarks on Protest

- The additional search fees were accompanied by the applicant's protest.
 No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

International application No
PCT/GB 95/01213

Patent document cited in search report	Publication date	Patent family member(s)		Publication date
WO-A-9401576	20-01-94	AU-B-	4667593	31-01-94
		CA-A-	2135193	20-01-94
		EP-A-	0654088	24-05-95
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WO-A-9315201	05-08-93	AU-B-	3482493	01-09-93
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		EP-A-	0624192	17-11-94
		JP-T-	7504813	01-06-95
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WO-A-9214748	03-09-92	EP-A-	0536350	14-04-93
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